CLEAN VERSION OF AMENDED CLAIMS

ENERATION OF RECOMBINANT ADENOVIRAL VECTOR SECH CENTER 1600/2900 Applicant: Beverly L. Davidson et al.

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- The cloning system of claim 16, wherein any or all open reading frames constituting E4 2. have been modified in the backbone plasmid.
- The cloning system of claim 2, wherein the modification is a substitution, insertion, or 3. deletion of one or more nucleotides.
- The cloning system of claim 16, wherein E3 has been modified in the backbone plasmid. 4.
- The cloning system of claim 4, wherein the modification is a substitution, insertion, or 5. deletion of one or more nucleotides.
- The cloning system of claim 4, wherein E3 has been modified to contain a multiple 6. cloning site.
- The cloning system of claim 4, wherein one or more genes required for Herpes Simplex 7. Virus (HSV) packaging and an HSV origin of replication have been placed within the E3 region.
- The cloning system of claim 16, further comprising in the backbone plasmid HSV 8. Amplicon sequences required for packaging and replication.
- The cloning system of claim 16, further comprising in the backbone plasmid one or more 10. sequences that allow for integration of sequences into cells after viral infection.
- A shuttle plasmid consisting essentially of Ad sequences from 0 to 1 and 9.2 to 16.1 map 11. units of an Ad genome.
 - The shuttle plasmid of claim 11, wherein PacI restriction endonuclease sites flank either 12. end of the Ad sequences.
 - The shuttle plasmid of claim 11, further comprising a multiple cloning site positioned 13. between 1 and 9.2 map units.
 - The shuttle plasmid of claim 11/, wherein the shuttle plasmid further comprises a 14. sequence encoding a gene of interest.
 - The shuttle plasmid of claim 11, further comprising a novel promoter, inducible promoter 15. or other sequence used to drive expression from a transgene.
 - A cloning system for generating recombinant adenovirus comprising: 16.
 - an Ad backbone plasmid consisting essentially of an Ad genome lacking map units 0 to 9.2, wherein the numbering of the map units starts with the lefthand ITR, and

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(b) a shuttle plasmid consisting essentially of Ad sequences from 0 to 1 and 9.2 to 16.1 map units of an Ad genome.

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17. A host cell comprising:

- (a) an Ad backbone plasmid consisting essentially of an Ad genome lacking map units 0 to 9.2, wherein the numbering of the map units starts with the lefthand ITR, and
- (b) a shuttle plasmid consisting essentially of Ad sequences from 0 to 1 and 9.2 to 16.1 map units of an Ad genome.
- 18. The host cell of claim 17, wherein the dell expresses E1 sequences necessary for supporting adenovirus replication.
- 19. A host cell of claim 18, wherein the kell is an animal cell.
- 20. A host cell of claim 17, wherein the cell expresses E1 sequences, pIX and E4 sequences required for amplification of viruses generated made with the Ad backbone lacking E1, E1 and pIX, or E1 and E4, respectively.
- 21. A host cell of 20, wherein the cell is an animal cell.
- 22. A method for rapidly producing recombinant adenovirus comprising contacting a host cell with
 - (a) an Ad backbone plasmid consisting essentially of an Ad genome lacking map units 0 to 9.2, wherein the numbering of the map units starts with the lefthand ITR, and
 - (b) a shuttle plasmid consisting essentially of Ad sequences from 0 to 1 and 9.2 to 16.1 map units of an Ad genome.
- 23. The method of claim 22, further comprising serially amplifying virus produced by the host cell.
- 24. The method of claim 23, further comprising detecting the presence of wild type virus.
- 25. The method of claim 22, wherein the shuttle plasmid further comprises a sequence encoding a gene of interest.